PATENT COOPERATION TREATY

INTERNATIONAL PRELIMINARY REPORT ON PATENTABILITY (Chapter II of the Patent Cooperation Treaty)

(PCT Article 36 and Rule 70)

13127PC2	FOR FURTHER ACTION	See Form PCT/IPEA/416
International application No. PCT/AU2005/000500	International filing date (day/month/y 6 April 2005	i of management
International Patent Classification (II	PC) or national classification and IPC	8 April 2004
Int. CL	12 0	
C12Q 1/68 (2006.01)		
Applicani		
THE STATE OF QUEENS.	LAND ACTING THROUGH ITS DEPAI	RTMENT OF HEALTH et al
	iminary examination report, established by thinsmitted to the applicant according to Article	s International Preliminary Examining
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Same was accommended by	ANNEXES, comprising:	
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International application No.
PCT/AU2005/000500

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		ini	mational search (under Rules 12.3(a) and 23.1 (b))	
		tury	lication of the international application (under Rule 1	2.4(a))
		inte	mational preliminary examination (Rules 55.2(a) and	(or \$5.3(a))
Ĵ	urnishe	ed to the	he elements of the international application, this repo receiving Office in response to an invitation under As of annexed to this report;	xt is based on (replacement sheets which have been ticle 14 are referred to in this report as "originally
] the	e interna	tional application as originally filed/furnished	
	X the	e descrip	tion:	
}			pages 1-29 as originally filed/furnished	
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	ì		table(s) related to the sequence listing (specify):	
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Claims

International application No.

NO

PCT/AU2005/000500

Box No. V	lox No. V Reasoned statement under Article 35(2) with regard to novelty, inventive step or industrial applicability citations and explanations supporting such statement			
i. Statement				
N	ovelty (N)	Claims 1-24	171	YES
		Claims		NO
101	ventive step (IS)	Claims 1-24		YES
		Claims		NO
înc	iustrial applicability (IA)	Claims 1-24		VES

2. Citations and explanations (Rule 70.7)

The following documents identified in the International Search Report have been considered for the purposes of this report:

- D1 Whiley, D. M. et al., 2004 (July), European Journal of Clinical and Microbial Infectious Diseases, 23: 705-710.
- D2 Ghustein, J. Z. et al., 1999, Molecular Diagnosis, 4: 233-239.

The applicant's invention resides in a method for the detection of *Netsseria gonorrhoeae* based on the detection of the porA pseudogene. The use of the porA pseudogene as a target for *N. gonorrhoeae* detection provides improved clinical sensitivity and specificity when compared to other PCR-based detection methods (p. 6 lines 22-28).

P Category Documents

D1 Whiley, D. M. et al., 2004, European Journal of Clinical and Microbial Infectious Diseases, 23: 705-710.

With regard to the document listed above, this is a document that was published prior to the international filing date but later than the priority date claimed, and which would otherwise be considered to be of particular relevance to the present application. Under the PCT, novelty is considered only in respect of documents published before the priority date. Should there be any issue with the priority date presently claimed this document may be relevant to the novelty and/or inventiveness of the invention claimed.

Novelty (N) and Inventive Step (IS)

D2 discloses a PCR-based simplex assay for the detection of N. meningitidis and N. genorrhoeae. In the method described primers directed to the parA gene are used to amplify the gene and the PCR product is then detected using an internal oligonucleotide probe for parA (p.234, Primer Synthesis for PCR and Liquid Hybridization-Gel Retardation Analysis of Amplification Products). While the method described discloses primers that enable the detection of both N. meningitidis and N. gonorrhoeae, there is no disclosure of primers which are not capable of hybridising to a porA nucleic acid of N. meningitidis as presently claimed. Therefore the subject matter of claims 1-24 is new and meets the requirements of Article 33(2) PCT with regard to novelty.

Continued in Supplemental Box

International application No. PCT/AU2005/000500

Supp	upplemental Box Relating to Sequence Listing	
Centi	ontinuation of Box No. 1, item 2:	
l. W	With regard to any nucleotide and/or amino acid sequence disclosed in the international application and claimed invention, this report was established on the basis of:	tecessary to the
à.	a. type of material	
	a sequence listing	
	X table(s) related to the sequence listing	
b.	b. format of material	
•	X on paper	
	in electronic form	
¢,		
	X contained in the international application as filed	
}	filed together with the international application in electronic form	
	furnished subsequently to this Authority for the purposes of search and/or examination	
	received by this Authority as an amendment* on	
2.	In addition, in the case that more than one version or copy of a sequence listing and/or table(s) relatiled or furnished, the required statements that the information in the subsequent or additional copi in the application as filed or does not go beyond the application as filed, as appropriate, were furni	es is identical to that
લ હતા	Additional comments:	
Oescrip	Figure 1 do not clearly identify the primers according to their SEQ ID NO. Based on the information pro- teription of Figures the claims have been searched on the assumption that the SEQ ID NOs correspond to ture 1 as set out below.	vided in the <u>Brief</u> he primers listed in
8	* NG-pap-1 = SEQ ID NO: 1	
×	* NG-pap 2 = SEQ ID NO: 2	
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International application No.

PCT/AU2005/000500

Supplemental Box

Continuation of Box V: Reasoned statement under Article 35(2) with regard to novelty, inventive step or industrial applicability

Furthermore, claims 1-24 meet the criteria set out in PCT Article 33(3) with regard to the requirement of Inventive Step. As discussed previously, the prior art discloses an assay that allows the detection of N. gonorrhoeae and N. maningitidis. Thus, the citation does not obviously suggest to the person skilled in the art that an assay based on the detection of the pseudogene por A would be capable distinguishing between N. gonorrhoeae and N. meningitidis. The method described in the prior art does not allow the discrimination of N. gonorrhoeae from N. meningitidis and there is no indication that sufficient variability in the por A gene of these two Neisseria species exists such that discrimination of the two species would be possible.

Contract.	Industrial Applicability (IA)
The same of the sa	The invention defined in the claims is considered to meet the requirements of Industrial Applicability under Article 33(4) of the PCT.
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CLAIMS

- 1. A method of determining whether an individual is or has been infected with Neisseria gonorrhoeae, said method including the step of using one or more oligonucleotides to detect said isolated por nucleic acid of Neisseria gonorrhoeae, if present in a biological sample obtained from said individual, a presence of said por nucleic acid indicating that said individual is or has been infected with Neisseria gonorrhoeae, wherein said one or more oligonucleotides are not capable of hybridizing to a por nucleic acid of Neisseria meningitidis sufficiently to enable detection of said por nucleic acid of Neisseria meningitidis if present in said biological sample.
- 2. The method of Claim 1, wherein said method includes the step of distinguishing said isolated por Anucleic acid of Neisseria gonorrhoeae from a por Anucleic of Neisseria memingitidis present in said biological sample.
- 3. The method of Claim 2, wherein said por Anucleic acid of Neisseria gonorrhoeae is distinguished from another Neisseria species other than N. meningitidis.
- 4. The method of Claim 1, including the step of subjecting the biological sample to mucleic acid sequence amplification under conditions which facilitate amplification of said isolated por A nucleic acid of Neisseria gonorrhoeae to produce an amplification product.
- 5. The method of Claim 4, wherein the amplification product corresponds to a fragment of a Neisseria gonorrhoeae porA pseudogene.
- 6. The method of Claim 5, wherein nucleic acid sequence amplification is performed under conditions which facilitate amplification of said isolated por Anucleic acid of Neisseria gonorrhoeae to a detectable level but which do not facilitate amplification of said por Anucleic of N. meningitidis to a detectable level.
- 7. The method of Claim 6, wherein nucleic acid sequence amplification is performed using one or more PCR primers having a nucleotide sequence selected from the group consisting of SEQ ID NO:1 and SEQ ID NO:2.
- The method of Claim 4, wherein said one or more oligonucleotides comprise a probe for detecting said amplification product by probe hybridization.

Amended Sheet IPEA/AU

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- 9. The method of Claim 8, wherein the probe is has a nucleotide sequence selected from the group consisting of SEQ ID NO:3; SEQ ID NO:4; SEQ ID NO:5; SEQ ID NO:6; SEQ ID NO:7; SEQ ID NO:8; SEQ ID NO:9.
- 10. The method of Claim 9, wherein the probe is has a nucleotide sequence selected from the group consisting of SBQ ID NO:3 and SEQ ID NO:4.
- 11. The method of Claim 8, wherein detection of said amplification product is performed using fluorescence resonance energy transfer (FRET).
- 12. A method of determining whether a human individual is or has been infected with Neisseria gonorrhoeae, , said method including the steps of:
- (i) subjecting a biological sample obtained from said human individual to nucleic acid sequence amplification using primers having respective nucleotide sequences according to SEQ ID NO:1 and SEQ ID NO:2, to produce a porA Neisseria gonorrhoeae amplification product from a Neisseria gonorrhoeae porA nucleic acid if present in said biological sample; and
- (ii) detecting said amplification product, if present, by probe hybridization and fluorescence resonance energy transfer (FRET) using oligonucleotides having respective nucleotide sequences according to SEQ ID NO:3 having a donor fluorophore and SEQ ID NO:4 having an acceptor fluorophore, whereby a presence of said porA amplification product indicates that said individual is or has been infected with Neisseria gonorrhoeae,
- 13. An ofigonucleotide which is capable of hybridizing to a porA nucleic acid of Neisseria gonorrhoeae sufficiently to enable detection of said porA nucleic acid, but which is not capable of hybridizing to a porA nucleic acid of Neisseria meningitidis sufficiently to enable detection of said porA nucleic acid of Neisseria meningitidis.
- 14. The oligonucleotide of Claim 12, wherein said oligonucleotide is not capable of hybridizing to a porA nucleic acid of another Neisseria species: other than N. meningitidis.
- 15. The oligonucleotide of Claim 14 having a nucleotide sequence selected from the group consisting of SEQ ID NO:3; SEQ ID NO:4; SEQ ID NO:5; SEQ ID NO:6; SEQ ID NO:7; SEQ ID NO:8; SEQ ID NO:9.

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- 16. The oligonucleotide of Claim 15 having a nucleotide sequence selected from the group consisting of SEQ ID NO:3 and SEQ ID NO:4.
- 17. A kit for detecting a porA nucleic acid of Neisseria gonorrhoeae, said kit comprising one or more oligonucleotides according to Claim 13 together with a DNA polymerase and/or one or more detection reagents.
- 18. The kit of Claim 17, wherein the one or more oligonucleotides have a nucleotide sequence selected from the group consisting of SEQ ID NO:3; SEQ ID NO:4; SEQ ID NO:5; SEQ ID NO:6; SEQ ID NO:7; SEQ ID NO:8; SEQ ID NO:9.
- 19. The kit of Claim 18, wherein the one or more oligonucleotides have a nucleotide sequence selected from the group consisting of SEQ ID NO:3 and SEQ ID NO:4.
- 20. The kit of Claim 17, further comprising one or more primers that facilitate amplification of an Neisseria genorrhoeae, purA nucleic acid.
- 21. The kit of Claim 20, wherein the one or more primers have a nucleotide sequence selected from the group consisting of SEQ ID NO:1 and SEQ ID NO:2.
- 22. A nucleic acid array comprising one or more ofigonucleotides according to Claim 13, immobilized, coupled, bound, impregnated or otherwise in communication with a substrate,
- 23. The nucleic acid array of Claim 22, wherein the one or more oligonucleotides have a nucleotide sequence selected from the group consisting of SEQ ID NO:3; SEQ ID NO:4; SEQ ID NO:5; SEQ ID NO:6; SEQ ID NO:7; SEQ ID NO:8; SEQ ID NO:9.
- 24. The nucleic acid array of Claim 23, wherein the one or more oligonucleotides have a nucleotide sequence selected from the group consisting of SEQ ID NO:3 and SEQ ID NO:4.